

**REMARKS**

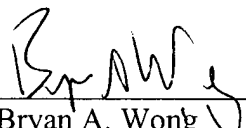
Applicants have received and reviewed an Office Communication dated October 2, 2002. By way of response, Applicants have amended the specification to include sequence identifiers in the appropriate places as requested by the Examiner.

Applicants respectfully request the Examiner to enter the above amendment. No new matter is introduced in the amendment.

Respectfully submitted,

MERCHANT & GOULD P.C.  
P.O. Box 2903  
Minneapolis, Minnesota 55402-0903  
(612) 332-5300

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Bryan A. Wong  
Reg. No. 50,836  
MDS:BAW:PSTkaw

**VERSION WITH MARKINGS TO SHOW CHANGES MADE**

**In the Specification**

Paragraphs beginning at page 6, line 28 have been amended as follows:

--Fig. 1 shows an alignment comparison of nucleotide sequences of HDGF2 (SEQ ID NO: 3) of the invention and mouse HDGF2 (SEQ ID NO: 9). The identical nucleotides are indicated by "|".

Fig. 2 shows an alignment comparison of amino acid sequences of HDGF2 (SEQ ID NO: 4) of the invention and mouse HDGF2 (SEQ ID NO:10). The identical and similar amino acids are indicated by "|" and ".", respectively.--

Paragraph beginning at page 6, line 33 has been amended as follows:

--In one embodiment, the cDNA sequence of HDGF2 was obtained as follows: human testis  $\lambda$ gt 11 cDNA library (Clontech) was used as a template and PCR was carried out with the synthetic forward primer A1 :5'-ACCGCTCGTC CGCCCGGCTT GAG-3' (SEQ ID NO: 1) and reverse primer B :5'-GATCCTAGAC ATGTATAAGT CTGCGC-3' (SEQ ID NO: 2). Target fragments of 1024bp were obtained. The sequencing of the PCR product gave the full length cDNA sequence shown in SEQ ID NO: 3.--